

## SEQUENCE LISTING

Vaisvila, Romualdus <110> Morgan, Richard D. Kucera, Rebecca B. Claus, Toby B. Raleigh, Elisabeth A. <120> Method For Cloning And Producing The MseI Restriction Endonuclease <130> NEB-181 <140> US 09/689,343 141> 2000-10-12 <160> <170> PatentIn version 3.1 <210> <211> 903 <212> DNA <213> Micrococcus sp. <220> <221> CDS <222> (1)..(900)<223> <400> 1 atg cet atc teg acc gtc tgg acg ceg gac gga gac gac etc atc gtg 48 Met Pro Ile Ser Thr Val Trp Thr Pro Asp Gly Asp Asp Leu Ile Val 10 96 gag gcg gac aac ctc gat ttc/att caa acg ctc ccc gac gcg agc ttc Glu Ala Asp Asn Leu Asp Phé Ile Gln Thr Leu Pro Asp Ala Ser Phe 25 cga atg atc tac atc gat /ccg ccg ttc aac aca ggg cga acg cag cgg 144 Arg Met Ile Tyr Ile Asp/Pro Pro Phe Asn Thr Gly Arg Thr Gln Arg ctt cag tcg ctc aag acg acc cgc tcg gtc aca ggg tcg cga gtc ggc 192 Leu Gln Ser Leu Lys /Thr Thr Arg Ser Val Thr Gly Ser Arg Val Gly 50 55 60 ttc aaa ggc cag acg tac gac acg gtc aag agc act ctg cac tcg tat 240 Phe Lys Gly Gln Thr Tyr Asp Thr Val Lys Ser Thr Leu His Ser Tyr 80 65 70 288 gac gac gct ttc acc gac tat tgg tcg ttc ctc gaa ccg cgt ctc ctg Asp Asp Ala Phé Thr Asp Tyr Trp Ser Phe Leu Glu Pro Arg Leu Leu 95 85 336 gag get tgg égg ttg etc acc ect gae gge geg etc tat ett eat etg

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| Glu Ala Trp Arg Leu Leu Thr Pro Asp Gly Ala Leu Tyr Leu His Leu 100 gat tac cgc gag gtt cac tac gcc aag gtc gtc ctc gac gcg atg ttc Asp Tyr Arg Glu Val His Tyr Ala Lys Val Val Leu Asp Ala Met Phe 115 gga cgc gaa agc ttc ctg aac gag ctg atc tgg gcg tac gac tac ggc Gly Arg Glu Ser Phe Leu Asn Glu Leu Ile Trp Ala Tyr Asp Tyr Gly 130 gcg cgc tcg aag agc aag tgg ccc acc aag cac gac aac atc ctc gtg Ala Arg Ser Lys Ser Lys Trp Pro Thr Lys His Asp Asn Ile Leu Val | 384<br>432<br>480 |
|---|-------------------|
| Asp Tyr Arg Glu Val His Tyr Ala Lys Val Val Leu Asp Ala Met Phe 115 120 125 125 125 125 125 125 125 125 125 125   | 432               |
| Gly Arg Glu Ser Phe Leu Asn Glu Leu Ile Trp Ala Tyr Asp Tyr Gly 130 135 140  gcg cgc tcg aag agc aag tgg ccc acc aag cac gac aac atc ctc gtg Ala Arg Ser Lys Ser Lys Trp Pro Thr Lys His Asp Asn Ile Leu Val  |                   |
| Ala Arg Ser Lys Ser Lys Trp Pro Thr Lys His Asp Asn Ile Leu Val   | 480               |
| 145 150 155 160   |                   |
| tat gtg aag gac ccg aac aac tac gtc tgg aac ggt cag gat gta gat Tyr Val Lys Asp Pro Asn Asn Tyr Val Trp Asn Gly Gln Asp Val Asp 165 170 175   | 528               |
| cgc gag ccc tac atg gcg ccc ggg ctc gtt aca ccc gag aag gta gcg<br>Arg Glu Pro Tyr Met Ala Pro Gly Leu Val Thr Pro Glu Lys Val Ala<br>180 185 190   | 576               |
| ctt ggc aag ctg ccc acc gac gtc tgg tgg cac aca atc gtt ccg cct<br>Leu Gly Lys Leu Pro Thr Asp Val Trp Trp His Thr Ile Val Pro Pro<br>195 200 205   | 624               |
| gcg agc aaa gag cgc acc ggg tac gcg aca cag aag ccg gtc ggc atc Ala Ser Lys Glu Arg Thr Gly Tyr Ala Thr Gln Lys Pro Val Gly Ile 210 215 220   | 672               |
| atc cgt cgc atg att cag gcg agc agc aat gaa ggc gac tgg gtt ctg  Ile Arg Arg Met Ile Gln Ala Ser Ser Asn Glu Gly Asp Trp Val Leu  235 230 235 240   | 720               |
| gat ttc ttc gct ggt agt ggg acg acc ggc gcc gcg gcc cgc cag ctc Asp Phe Phe Ala Gly Ser Gly Thr Thr Gly Ala Ala Ala Arg Gln Leu 245 250 255   | 768               |
| gga cgc cgt ttt gtg ctc gta gac gtc aac cca gaa gca atc gcg gta<br>Gly Arg Arg Phe Val Leu Val Asp Val Asn Pro Glu Ala Ile Ala Val<br>260 . 265 . 270   | 816               |
| atg gca aaa cgg ttg gat gac ggg gca ttg gac acc agc gtg acg atc Met Ala Lys Arg Leu Asp Asp Gly Ala Leu Asp Thr Ser Val Thr Ile 275 280 285   | 864               |
| gtg cag act ccc cag agt gac cca cga acc gac gga tga Val Gln Thr Pro Gln Ser Asp Pro Arg Thr Asp Gly 290 295 300   | 903               |

<sup>&</sup>lt;210> 2

<sup>&</sup>lt;211> 300 <212> PRT <213> Micrococcus sp.

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Glu Ala Asp Asn Leu Asp Phe Ile Gln Thr Leu Pro Asp Ala Ser Phe
20 25 30

Arg Met Ile Tyr Ile Asp Pro Pro Phe Asn Thr Gly Arg Thr Gln Arg 35 40 45

Leu Gln Ser Leu Lys Thr Thr Arg Ser Val Thr Gly Ser Arg Val Gly 50 55 60

Phe Lys Gly Gln Thr Tyr Asp Thr Val Lys Ser Thr Leu His Ser Tyr 65 70 75 80

Asp Asp Ala Phe Thr Asp Tyr Trp Ser Phe Leu Glu Pro Arg Leu Leu 85 90 95

Glu Ala Trp Arg Leu Leu Thr Pro Asp Gly Ala Leu Tyr Leu His Leu
100 105 110

Asp Tyr Arg Glu Val His Tyr Ala Lys Val Val Leu Asp Ala Met Phe 115 120 125

Gly Arg Glu Ser Phe Leu Asn Glu Leu Ile Trp Ala Tyr Asp Tyr Gly 130 135 140

Ala Arg Ser Lys Ser Lys Trp Pro Thr Lys His Asp Asn Ile Leu Val 145 150 155 160

Tyr Val Lys Asp Pro Asn Asn Tyr Val Trp Asn Gly Gln Asp Val Asp 165 170 175

Arg Glu Pro Tyr Met Ala Pro Gly Leu Val Thr Pro Glu Lys Val Ala 180 185 190

Leu Gly Lys Leu Pro Thr Asp Val Trp Trp His Thr Ile Val Pro Pro 195 200 205

Ala Ser Lys Glu Arg Thr Gly Tyr Ala Thr Gln Lys Pro Val Gly Ile 210 215 220

Ile Arg Arg Met Ile Gln Ala Ser Ser Asn Glu Gly Asp Trp Val Leu 225 230 235 240

Asp Phe Phe Ala Gly Ser Gly Thr Thr Gly Ala Ala Ala Arg Gln Leu 245 250 255

Gly Arg Arg Phe Val Leu Val Asp Val Asn Pro Glu Ala Ile Ala Val 260 265 270

Met Ala Lys Arg Leu Asp Asp Gly Ala Leu Asp Thr Ser Val Thr Ile 275 280 285

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                        295
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       Xaa = any amino acid
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      misc_feature
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                                                                        48
Met Pro Thr Leu Asp Trp Pro Gly Lys Gln Leu Ser Phe Pro Pro Ala
                                                         15
                5
                                                                        96
acc tcc ttg cat ctg gag agt gtg gtc act gag gga gcg gag tca ccg
Thr Ser Leu His Leu Glu Ser Val Val Thr Glu Gly Ala Glu Ser Pro
                                                                       144
cct aat cgt ctg att tgg gcg gac aac ctg ccg cta atg gta gat ttg
Pro Asn Arg Leu Ile Trp Ala Asp Asn Leu Pro Leu Met Val Asp Leu
                                                                       192
ttg gcc gaa tat gaa ggg aaa atc gat ctg atc tac gcc gat ccc cct
Leu Ala Glu Tyr Glu Gly Lys Ile Asp Leu Ile Tyr Ala Asp Pro Pro
    50
                                                                       240
ttt ttt acg gat cgt act tat gcg gcg cga att ggt cat ggg gag gat
Phe Phe Thr Asp Arg Thr Tyr Ala Ala Arg Ile Gly His Gly Glu Asp
                    70
                                         75
                                                              80
65
teg egt egt eca caa ace tgg eag ett gea gaa gga tat acg gae gag
                                                                       288
Ser Arg Arg Pro Gln Thr Trp Gln Leu Ala Glu Gly Tyr Thr Asp Glu
                85
tgg aag gat tta gat gaa tac ctg gac ttc ctt tat cca cgc ctg gta
                                                                       336
Trp Lys Asp Leu Asp Glu Tyr Leu Asp Phe Leu Tyr Pro Arg Leu Val
            100
                                 105
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|                       |   |   | , |   |   |   |   |      |   |       |   |      |
|-----------------------|---|---|---|---|---|---|---|------|---|-------|---|------|
| atg<br>Met            |   |   |   |   |   |   |   |      |   |       |   | 384  |
| tgg<br>Trp<br>130     |   |   |   |   |   |   |   |      |   |       |   | 432  |
| cga<br>Arg            |   |   |   |   |   |   |   |      |   |       |   | 480  |
| gcc<br>Ala            |   |   |   |   |   |   |   |      |   |       |   | 528  |
| gtg<br>Val            |   |   | _ |   |   |   |   |      | _ | <br>_ | - | 576  |
| tac<br>Tyr            |   | _ | _ |   |   | _ |   |      |   |       |   | 624  |
| <br>ttt<br>Phe<br>210 |   |   |   |   |   |   |   |      |   |       |   | 672  |
| tgg<br>Trp            |   |   |   |   |   |   |   |      |   |       |   | 720  |
| ccg<br>Pro            |   |   | _ |   |   | _ | _ | <br> |   | _     |   | 768  |
| tcg<br>Ser            |   |   |   |   |   |   |   |      |   |       |   | 816  |
| acc<br>Thr            | _ |   |   | - | _ | - |   |      |   |       |   | 864  |
| gca<br>Ala<br>290     |   |   |   |   |   |   |   |      |   |       |   | 912  |
| <br>gga<br>Gly        | - | _ |   |   |   |   |   |      |   |       |   | 960  |
| cag<br>Gln            |   |   |   |   |   |   |   |      |   |       |   | 1008 |

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|--------------------------|----------------------|-----------------------|-----------|----------|-------|-----------|-----------|-----------|-----------|-----|------------|-----------|------------|-----------|-----|------|
|                          |                      |                       |           | •        |       |           |           |           |           |     |            |           |            |           |     |      |
|                          |                      | caa<br>Gln            |           |          |       |           |           |           |           |     |            |           |            |           |     | 1056 |
|                          |                      | ggc<br>Gly<br>355     |           |          |       |           |           |           |           |     |            |           |            |           |     | 1104 |
|                          |                      | gag<br>Glu            |           |          |       |           |           |           |           |     |            |           |            |           |     | 1152 |
|                          | Cys                  | gta<br>Val            |           |          |       |           |           |           |           |     |            |           |            |           |     | 1200 |
|                          |                      | gcc<br>Ala            |           |          |       |           |           |           |           |     | tga        |           |            |           |     | 1236 |
| <21<br><21               | 0><br>1><br>2><br>3> | 411                   | own       |          |       |           |           |           |           |     |            |           |            |           |     |      |
| <22<br><22               |                      | Envi:                 | ronme     | enta:    | l DNA | A         |           |           |           |     |            |           |            |           |     |      |
| <22<br><22<br><22<br><22 | 1><br>2>             | misc<br>(198<br>Xaa = | ) (       | 198)     | ino a | acid      |           |           |           |     |            |           |            |           |     |      |
| <22                      | 1><br>2>             | misc<br>(594)<br>N= G | ) (       | 594)     | r T   |           |           |           |           |     |            |           |            |           |     |      |
| <40                      | 0>                   | 4                     |           |          |       |           |           |           |           |     |            |           |            |           |     |      |
| Met<br>1                 | Pro                  | Thr                   | Leu       | Asp<br>5 | Trp   | Pro       | Gly       | Lys       | Gln<br>10 | Leu | Ser        | Phe       | Pro        | Pro<br>15 | Ala |      |
| Thr                      | Ser                  | Leu                   | His<br>20 | Leu      | Glu   | Ser       | Val       | Val<br>25 | Thr       | Glu | Gly        | Ala       | Glu<br>30  | Ser       | Pro |      |
| Pro                      | Asn                  | Arg<br>35             | Leu       | Ile      | Trp   | Ala       | Asp<br>40 | Asn       | Leu       | Pro | Leu        | Met<br>45 | Val        | Asp       | Leu |      |
| Leu                      | Ala<br>50            | Glu                   | Tyr       | Glu      | Gly   | Lys<br>55 | Ile       | Asp       | Leu       | Ile | Tyr<br>60  | Ala       | Asp        | Pro       | Pro |      |
|                          |                      |                       |           |          |       | _         |           | Ala       |           | -1- | <b>6</b> 3 | ***       | <b>~</b> 1 | <b>01</b> | 3   |      |

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Ser Arg Arg Pro Gln Thr Trp Gln Leu Ala Glu Gly Tyr Thr Asp Glu Trp Lys Asp Leu Asp Glu Tyr Leu Asp Phe Leu Tyr Pro Arg Leu Val 105 Leu Met Tyr Arg Leu Leu Ala Pro His Gly Thr Leu Tyr Leu His Leu 120 Asp Trp His Ala Asn Ala Tyr Val Arg Val Leu Leu Asp Glu Ile Phe 135 Gly Arg Gln Arg Phe Leu Asn Glu Ile Val Trp Ile Tyr His Gly Pro 145 150 Ser Ala Ile Arg Arg Ala Phe Lys Arg Lys His Asp Thr Ile Leu Val 170 Tyr Val Lys Gly Glu Asn Tyr Thr Phe Asn Ala Asp Ala Val Arg Gln Pro Tyr His Pro Ser Xaa His Lys Thr Phe Ala Ser Ser Pro Lys Ala 195 200 Gly Phe Gly Lys Val Pro Asp Leu Gln Arg Gly Lys Val Pro Glu Asp Trp Trp Tyr Phe Pro Val Val Ala Arg Leu His Arg Glu Arg Ser Gly 230 235 Tyr Pro Thr Gln Lys Pro Gln Ala Leu Leu Glu Arg Ile Leu Leu Ala 245 250 Ser Ser Asn Ala Gly Asp Leu Val Ala Asp Phe Phe Cys Gly Ser Gly 265 Thr Thr Ala Val Val Ala Ala Arg Leu Gly Arg Arg Phe Leu Val Asn 275 280 Asp Ala Ser Trp Arg Ala Val His Val Thr Arg Thr Arg Leu Leu Arg 295 Glu Gly Val Ser Phe Thr Phe Glu Arg Gln Glu Thr Phe Thr Leu Pro 310 315 Ile Gln Pro Leu Pro Pro Asp Trp Leu Ile Ile Ala Glu Glu Gln Ile 330 325 Arg Leu Gln Ala Pro Phe Leu Val Asp Phe Trp Glu Val Asp Asp Gln 345 Trp Asp Gly Lys Ile Phe Arg Ser Arg His Gln Gly Leu Arg Ser Arg 355

Leu Gln Glu Gln Ala Pro Leu Ser Leu Pro Leu Thr Gly Asn Gly Leu

|            |      |     | •   |            |            |     |     |     |            |            |     |     |     |     |            |
|------------|------|-----|-----|------------|------------|-----|-----|-----|------------|------------|-----|-----|-----|-----|------------|
|            | 370  |     |     |            |            | 375 |     |     |            |            | 380 |     |     |     |            |
| Leu<br>385 | Cys  | Val | Arg | Val        | Val<br>390 | Ser | Arg | Glu | Gly        | Glu<br>395 | Tyr | Tyr | Glu | Phe | Thr<br>400 |
| Gly        | Arg  | Ala | Asp | Ser<br>405 | Pro        | His | Pro | Val | Ser<br>410 | Phe        |     |     |     |     |            |
| -21(       | ۱۰ ۱ | =   |     |            |            |     |     |     |            |            |     |     |     |     |            |

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tac cgc gag gtg cac tac tgt aag gtc ctt ctt gac ggc atc ttc ggt 384
Tyr Arg Glu Val His Tyr Cys Lys Val Leu Leu Asp Gly Ile Phe Gly
115 120 125

|     | -            |      | gcc<br>Ala        |       |       |       | - |  |  |  |  | 4 | 132 |  |
|-----|--------------|------|-------------------|-------|-------|-------|---|--|--|--|--|---|-----|--|
|     |              |      | aag<br>Lys        |       |       |       |   |  |  |  |  | 4 | 80  |  |
|     |              |      | act<br>Thr        |       |       |       |   |  |  |  |  | 5 | 528 |  |
|     |              |      | tac<br>Tyr        |       |       |       |   |  |  |  |  | 5 |     |  |
| · · |              |      | ctg<br>Leu<br>195 |       |       |       |   |  |  |  |  | 6 | 524 |  |
|     |              |      | gag<br>Glu        | _     |       |       |   |  |  |  |  | 6 | 72  |  |
|     |              | -    | att<br>Ile        |       |       |       |   |  |  |  |  | 7 | 220 |  |
|     |              |      | gcc<br>Ala        |       |       |       |   |  |  |  |  | 7 | 68  |  |
|     |              |      | ttc<br>Phe        |       |       |       |   |  |  |  |  | 8 | 16  |  |
|     |              |      | cgc<br>Arg<br>275 |       |       |       |   |  |  |  |  | 8 | 64  |  |
|     | Pro          |      | ccg<br>Pro        |       |       |       |   |  |  |  |  | 9 | 12  |  |
|     |              |      | ggg<br>Gly        | taa   |       |       |   |  |  |  |  | 9 | 24  |  |
|     | <210<br><211 | .> 3 | 307               |       |       |       |   |  |  |  |  |   |     |  |
|     | <212<br><213 |      | PRT<br>Jnkno      | own   |       |       |   |  |  |  |  |   |     |  |
|     | <220<br><223 |      | Envir             | conme | ental | l dna | Ą |  |  |  |  |   |     |  |
|     |              |      |                   |       |       |       |   |  |  |  |  |   |     |  |

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## <400> 6

- Ala Asp Asn Met Glu Val Leu Arg Gly Leu Pro Ala Ala Ser Val Asp 20 25 30
- Leu Ile Tyr Ile Asp Pro Pro Phe Asn Thr Gly Lys Val Gln Glu Arg
  35 40 45
- Thr Gln Leu Lys Thr Val Arg Ser Glu Trp Gly Asp Arg Val Gly Phe 50 55 60
- Gln Gly Arg Arg Tyr Glu Ser Ile Val Val Gly Lys Lys Arg Phe Thr 65 70 75 80
- Asp Phe Phe Asp Asp Tyr Leu Ala Phe Leu Glu Pro Arg Leu Val Glu 85 90 95
- Ala His Arg Val Leu Ala Pro His Gly Cys Leu Tyr Phe His Val Asp 100 105 110
- Tyr Arg Glu Val His Tyr Cys Lys Val Leu Leu Asp Gly Ile Phe Gly
  115 120 125
- Arg Glu Ala Phe Leu Asn Glu Ile Ile Trp Ala Tyr Asp Tyr Gly Gly
  130 135 140
- Arg Pro Lys Asp Arg Trp Pro Pro Lys His Asp Asn Ile Leu Leu Tyr 145 150 155 160
- Ala Lys Thr Pro Gly Arg His Val Phe Asn Ala Asp Glu Ile Glu Arg 165 170 175
- Ile Pro Tyr Met Ala Pro Gly Leu Val Gly Pro Glu Lys Ala Ala Arg 180 185 190
- Gly Lys Leu Pro Thr Asp Thr Trp Trp His Thr Ile Val Pro Thr Ser 195 200 205
- Gly Ser Glu Lys Thr Gly Tyr Pro Thr Gln Lys Pro Leu Gly Ile Leu 210 215 220
- Arg Arg Ile Val Gln Ala Ser Ser His Pro Gly Ala Val Leu Asp 225 230 235 240
- Phe Phe Ala Gly Ser Gly Thr Thr Gly Val Ala Ala Phe Glu Leu Gly 245 250 255
- Arg Arg Phe Ile Leu Val Asp Asn His Pro Glu Ala Leu Gln Val Met 260 265 270
- Ala Arg Arg Phe Asp Gly Ile Glu Gly Ile Glu Trp Val Gly Phe Asp 275 280 285

Pro Thr Pro Tyr Gln Lys Gly Ala Lys Gln Arg Arg Ser Cys Pro Ala Pro Thr Gly 305 <210> 7 <211> 561 <212> DNA <213> Micrococcus sp. <220> <221> CDS <222> (1)..(558)<223> <400> 7 gtg acc cac gaa ccg acg gat gat ccc gat ttc ata gtg atg gcc gcg 48 Val Thr His Glu Pro Thr Asp Asp Pro Asp Phe Ile Val Met Ala Ala 10 age geg geg aac ete get gat egg tae gta geg agt gaa gae gee eee 96 Ser Ala Ala Asn Leu Ala Asp Arg Tyr Val Ala Ser Glu Asp Asp Pro tgg gtc ggc agc ccg ttc gag tgg atc ctt cgc gtt cca tcc aga acg 144 Trp Val Gly Ser Pro Phe Glu Trp Ile Leu Arg Val Pro Ser Arg Thr 35 192 aag ggc gcg gtc ggt gag ctg ctc gtg agc gaa tgg gct aat gcc aaa Lys Gly Ala Val Gly Glu Leu Leu Val Ser Glu Trp Ala Asn Ala Lys 50 55 ggc ctc cgt gtg aag agg tcg ggg tcc agc gat gcg gac cgc gtg atc 240 Gly Leu Arg Val Lys Arg Ser Gly Ser Ser Asp Ala Asp Arg Val Ile 65 70 288 aac ggg cat cgc atc gag atc aag atg tcg act ttg tgg aag tcc ggc Asn Gly His Arg Ile Glu Ile Lys Met Ser Thr Leu Trp Lys Ser Gly 85 90 95 336 ggc ttc aag ttt cag cag atc cgg gat cag gag tac gac ttt tgc ctc Gly Phe Lys Phe Gln Gln Ile Arg Asp Gln Glu Tyr Asp Phe Cys Leu 100 105 384 tgc ctt ggg atc agc ccg ttc gaa gtg cac gcg tgg ctg ctc aaa Cys Leu Gly Ile Ser Pro Phe Glu Val His Ala Trp Leu Leu Pro Lys 120 432 gac cta ttg ctt gag tac gtg att ggt cac atg ggt cag cac acc ggc Asp Leu Leu Glu Tyr Val Ile Gly His Met Gly Gln His Thr Gly

130

135

|  | •                  | ·                |                   |                   |                |            |
|--|--------------------|------------------|-------------------|-------------------|----------------|------------|
| gcg agc ggg<br>Ala Ser Gly<br>145                |                    |                  | Leu Gly Pl        |                   |                |            |
| tat gac tgg<br>Tyr Asp Trp                       |                    |                  |                   |                   |                |            |
| ctc ctc ctc<br>Leu Leu Leu                       |                    | Pro Gly P        |                   | ga                |                | 561        |
| <210> 8<br><211> 186<br><212> PRT<br><213> Micro | ococcus sp.        |                  |                   |                   |                |            |
| <400> 8  |                    |                  |                   |                   |                |            |
| Val Thr His<br>1                                 | Glu Pro Thr<br>5   | Asp Asp P        | Pro Asp Pl<br>10  | he Ile Val        | Met Ala<br>15  | Ala        |
| Ser Ala Ala                                      | Asn Leu Ala<br>20  |                  | Tyr Val A<br>25   | la Ser Glu        | Asp Asp<br>30  | Pro        |
| Trp Val Gly<br>35                                | Ser Pro Phe        | Glu Trp I<br>40  | Ile Leu A         | rg Val Pro<br>45  | Ser Arg        | Thr        |
| Lys Gly Ala<br>50                                | Val Gly Glu        | Leu Leu V<br>55  | /al Ser G         | lu Trp Ala<br>60  | Asn Ala        | Lys        |
| Gly Leu Arg<br>65                                | Val Lys Arg<br>70  | Ser Gly S        | Ser Ser As<br>7!  |                   | Arg Val        | Ile<br>80  |
| Asn Gly His                                      | Arg Ile Glu<br>85  | Ile Lys M        | Met Ser Tl<br>90  | hr Leu Trp        | Lys Ser<br>95  | Gly        |
| Gly Phe Lys                                      | Phe Gln Gln<br>100 |                  | Asp Gln G<br>105  | lu Tyr Asp        | Phe Cys<br>110 | Leu        |
| Cys Leu Gly<br>115                               | Ile Ser Pro        | Phe Glu V<br>120 | /al His A         | la Trp Leu<br>125 | Leu Pro        | Lys        |
| Asp Leu Leu<br>130                               | Leu Glu Tyr        | Val Ile G<br>135 | Gly His Me        | et Gly Gln<br>140 | His Thr        | Gly        |
| Ala Ser Gly<br>145                               | Ser Asp Thr<br>150 | Ala Trp L        |                   | he Pro Ala<br>55  | Asp Glu        | Pro<br>160 |
| Tyr Asp Trp                                      | Met Arg Pro<br>165 | Phe Gly G        | Gly Arg Le<br>170 | eu Gly His        | Val Glu<br>175 | Asp        |
| Leu Leu Leu                                      | Ala Ala Gly<br>180 |                  | Pro Tyr<br>185    |                   |                |            |

<211> 413 Escherichia coli <400> 9 accggtgatt ggacattgcc gaaatcaggc tgtctctcac tatttgacgc actggctgga 60 120 ctatccacat ctaccttatt cccccgaata acgagatccc ttccagcacc gggcaattgc 180 ccggtttttt ttgcgttgaa tttgtcattt tgtgccgtgg tgtttaaacc gcacagaata aattgtcgtg atttcacctt taaaataaaa ttaaaagaga aaaaaattct ctgtggaagg 240 gctatgttag ataaaattga ccgtaagctg ctggccttac tgcagcagga ttgcaccctc 300 tctttgcagg cactggctga agccgttaat ctgacaacca ccccttgctg gaagcgcctg 360 413 aaacggctgg aggacgacgg tatccttatc ggcaaagtcg ccctgctgga tcc

Chron

<210>